RAW SEQUENCE LISTING PATENT APPLICATION US/09/242,977

1632

DATE: 02/04/2000 TIME: 02:02:36

INPUT SET: S34634.raw

This Raw Listing contains the General Information Section and up to the first 5 pages.

1 2			SEQUENCE LISTING	ENTERED
3	(1)	General Information:		
5 6 7	(i) APPLICANT: Wilson, James Fisher, Krish		
8 9 10	(ii) TITLE OF INVENTION: Meth Virus-Directed Gene T	od for Recombinant Adeno herapy	-Associated
11 12	(iii) NUMBER OF SEQUENCES: 4		
13 14 15 16 17 18 19 20	(iv) CORRESPONDENCE ADDRESS: (A) ADDRESSEE: Howson a (B) STREET: Spring House (C) CITY: Spring House (D) STATE: Pennsylvania (E) COUNTRY: USA (F) ZIP: 19477	nd Howson e Corporate Cntr, PO Box	457
21 22 23 24 25 26	(v)	COMPUTER READABLE FORM: (A) MEDIUM TYPE: Floppy (B) COMPUTER: IBM PC cor (C) OPERATING SYSTEM: PC (D) SOFTWARE: Patentin F	mpatible	30
27 28 29 30 31	(vi)	CURRENT APPLICATION DATA: (A) APPLICATION NUMBER: (B) FILING DATE: (C) CLASSIFICATION:	us	
32 33 34 35	(vii)	PRIOR APPLICATION DATA: (A) APPLICATION NUMBER: (B) FILING DATE: 06-SEP-	US 08/708,188 1996	
36 37 38 39	(vii)	PRIOR APPLICATION DATA: (A) APPLICATION NUMBER: (B) FILING DATE: 10-OCT-	US 08/729,061 1996	
40 41 42 43 44	(viii)	ATTORNEY/AGENT INFORMATIO (A) NAME: Kodroff, Cathy (B) REGISTRATION NUMBER: (C) REFERENCE/DOCKET NUM	A. 33,980	
45 46	(ix)	TELECOMMUNICATION INFORMATION TELEPHONE: 215-540-9	FION: 200	

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DATE: 02/04/2000 TIME: 02:02:36

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47 48 49 50	(B) TELEFAX: 215-540-5818	INPUT SET: S34634.raw							
51 52	(2) INFORMATION FOR SEQ ID NO:1:								
53	(i) SEQUENCE CHARACTERISTICS:								
54	(A) LENGTH: 10398 base pairs								
55 56	(B) TYPE: nucleic acid								
57	(c) pid-indepliess: double								
58	(D) TOPOLOGY: unknown								
59 60	(ii) MOLECULE TYPE: cDNA								
61									
62									
63 64	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:1:								
65	GAATTCGCTA GCATCATCAA TAATATACCT TATTTTGGAT TGAAGCCAAT								
66			60						
67 68	GGGGGTGGAG TTTGTGACGT GGCGCGGGC GTGGGAACGG GGCGGGTGAC	GTAGTAGTGT	120						
69 70	GGCGGAAGTG TGATGTTGCA AGTGTGGCGG AACACATGTA AGCGACGGAT	GTGGCAAAAG	180						
71 72	TGACGTTTTT GGTGTGCGCC GGTGTACACA GGAAGTGACA ATTTTCGCGC	GGTTTTAGGC	240						
73	GGATGTTGTA GTAAATTTGG GCGTAACCGA GTAAGATTTG GCCATTTTCG	CGGGAAAACT	300						
74 75	GAATAAGAGG AAGTGAAATC TGAATAATTT TGTGTTACTC ATAGCGCGTA		360						
76 77	AGGGAGATCT GCTGCGCGCT CGCTCGCTCA CTGAGGCCGC CCGGGCAAAG								
78 79	GGGCGACCTT TGGTCGCCCG GCCTCAGTGA GCGAGCGAGC GCGCAGAGAG		420						
80			480						
81 82	ACTCCATCAC TAGGGGTTCC TTGTAGTTAA TGATTAACCC GCCATGCTAC		540						
83 84	TTCGAGCTTG CATGCCTGCA GGTCGTTACA TAACTTACGG TAAATGGCCC	GCCTGGCTGA	600						
85 86	CCGCCCAACG ACCCCCGCCC ATTGACGTCA ATAATGACGT ATGTTCCCAT	AGTAACGCCA	660						
87 88	ATAGGGACTT TCCATTGACG TCAATGGGTG GAGTATTTAC GGTAAACTGC	CCACTTGGCA	720						
89 90	GTACATCAAG TGTATCATAT GCCAAGTACG CCCCCTATTG ACGTCAATGA	CGGTAAATGG	780						
91	CCCGCCTGGC ATTATGCCCA GTACATGACC TTATGGGACT TTCCTACTTG	GCAGTACATC	840						
92 93	TACGTATTAG TCATCGCTAT TACCATGGTG ATGCGGTTTT GGCAGTACAT		900						
94 95	GGATAGCGGT TTGACTCACG GGGATTTCCA AGTCTCCACC CCATTGACGT								
96 97			960						
98	TTGTTTTGGC ACCAAAATCA ACGGGACTTT CCAAAATGTC GTAACAACTC		1020						
99	ACGCAAATGG GCGGTAGGCG TGTACGGTGG GAGGTCTATA TAAGCAGAGC	TCGTTTAGTG	1080						

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DATE: 02/04/2000 TIME: 02:02:37

INPUT SET: S34634.raw AACCGTCAGA TCGCCTGGAG ACGCCATCCA CGCTGTTTTG ACCTCCATAG AAGACACCGG GACCGATCCA GCCTCCGGAC TCTAGAGGAT CCGGTACTCG AGGAACTGAA AAACCAGAAA GTTAACTGGT AAGTTTAGTC TTTTTGTCTT TTATTTCAGG TCCCGGATCC GGTGGTGGTG CAAATCAAAG AACTGCTCCT CAGTGGATGT TGCCTTTACT TCTAGGCCTG TACGGAAGTG TTACTTCTGC TCTAAAAGCT GCGGAATTGT ACCCGCGGCC GCAATTCCCG GGGATCGAAA GAGCCTGCTA AAGCAAAAAA GAAGTCACCA TGTCGTTTAC TTTGACCAAC AAGAACGTGA TTTTCGTTGC CGGTCTGGGA GGCATTGGTC TGGACACCAG CAAGGAGCTG CTCAAGCGCG ATCCCGTCGT TTTACAACGT CGTGACTGGG AAAACCCTGG CGTTACCCAA CTTAATCGCC TTGCAGCACA TCCCCCTTTC GCCAGCTGGC GTAATAGCGA AGAGGCCCGC ACCGATCGCC CTTCCCAACA GTTGCGCAGC CTGAATGGCG AATGGCGCTT TGCCTGGTTT CCGGCACCAG AAGCGGTGCC GGAAAGCTGG CTGGAGTGCG ATCTTCCTGA GGCCGATACT GTCGTCGTCC CCTCAAACTG GCAGATGCAC GGTTACGATG CGCCCATCTA CACCAACGTA ACCTATCCCA TTACGGTCAA TCCGCCGTTT GTTCCCACGG AGAATCCGAC GGGTTGTTAC TCGCTCACAT TTAATGTTGA TGAAAGCTGG CTACAGGAAG GCCAGACGCG AATTATTTTT GATGGCGTTA ACTCGGCGTT TCATCTGTGG TGCAACGGGC GCTGGGTCGG TTACGGCCAG GACAGTCGTT TGCCGTCTGA ATTTGACCTG AGCGCATTTT TACGCGCCGG AGAAAACCGC CTCGCGGTGA TGGTGCTGCG TTGGAGTGAC GGCAGTTATC TGGAAGATCA GGATATGTGG CGGATGAGCG GCATTTTCCG TGACGTCTCG TTGCTGCATA AACCGACTAC ACAAATCAGC GATTTCCATG TTGCCACTCG CTTTAATGAT GATTTCAGCC GCGCTGTACT GGAGGCTGAA GTTCAGATGT GCGGCGAGTT GCGTGACTAC CTACGGGTAA CAGTTTCTTT ATGGCAGGGT GAAACGCAGG TCGCCAGCGG CACCGCGCCT TTCGGCGGTG AAATTATCGA TGAGCGTGGT GGTTATGCCG ATCGCGTCAC ACTACGTCTG AACGTCGAAA ACCCGAAACT GTGGAGCGCC GAAATCCCGA ATCTCTATCG TGCGGTGGTT GAACTGCACA CCGCCGACGG CACGCTGATT GAAGCAGAAG CCTGCGATGT CGGTTTCCGC GAGGTGCGGA TTGAAAATGG TCTGCTGCTG CTGAACGGCA TGGATGAGCA GACGATGGTG CAGGATATCC TGCTGATGAA GCAGAACAAC TTTAACGCCG

RAW SEQUENCE LISTING PATENT APPLICATION US/09/242,977

DATE: 02/04/2000 TIME: 02:02:37

153	INPUT SET: S340	534.raw
154	TGTGGTACAC COMPAGAAAA	2700
155 156	TGTATGTGGT GGATGAAGCC AATATTGAAA CCCACGGCAT GGTGCCAATG AATCGTCTGA	2760
157 158	CCGATGATCC GCGCTGGCTA CCGGCGATGA GCGAACGCGT AACGCGAATG GTGCAGCGCG	2820
159 160	ATCGTAATCA CCCGAGTGTG ATCATCTGGT CGCTGGGGAA TGAATCAGGC CACGGCGCTA	2880
161 162	ATCACGACGC GCTGTATCGC TGGATCAAAT CTGTCGATCC TTCCCGCCCG GTGCAGTATG	2940
163 164	AAGGCGGCGG AGCCGACACC ACGGCCACCG ATATTATTTG CCCGATGTAC GCGCGCGTGG	3000
165 166	ATGAAGACCA GCCCTTCCCG GCTGTGCCGA AATGGTCCAT CAAAAAATGG CTTTCGCTAC	3060
167 168	CTGGAGAGAC GCGCCCGCTG ATCCTTTGCG AATACGCCCA CGCGATGGGT AACAGTCTTG	3120
169 170	GCGGTTTCGC TAAATACTGG CAGGCGTTTC GTCAGTATCC CCGTTTACAG GGCGGCTTCG	3180
171 172	TCTGGGACTG GGTGGATCAG TCGCTGATTA AATATGATGA AAACGGCAAC CCGTGGTCGG	3240
173 174	CTTACGGCGG TGATTTTGGC GATACGCCGA ACGATCGCCA GTTCTGTATG AACGGTCTGG	3300
175 176	TCTTTGCCGA CCGCACGCCG CATCCAGCGC TGACGGAAGC AAAACACCAG CAGCAGTTTT	3360
177 178	TCCAGTTCCG TTTATCCGGG CAAACCATCG AAGTGACCAG CGAATACCTG TTCCGTCATA	3420
179 180	GCGATAACGA GCTCCTGCAC TGGATGGTGG CGCTGGATGG TAAGCCGCTG GCAAGCGGTG	3480
181 182	AAGTGCCTCT GGATGTCGCT CCACAAGGTA AACAGTTGAT TGAACTGCCT GAACTACCGC	3540
183 184	AGCCGGAGAG CGCCGGGCAA CTCTGGCTCA CAGTACGCGT AGTGCAACCG AACGCGACCG	3600
185 186	CATGGTCAGA AGCCGGGCAC ATCAGCGCCT GGCAGCAGTG GCGTCTGGCG GAAAACCTCA	3660
187 188	GTGTGACGCT CCCCGCCGCG TCCCACGCCA TCCCGCATCT GACCACCAGC GAAATGGATT	3720
189 190	TTTGCATCGA GCTGGGTAAT AAGCGTTGGC AATTTAACCG CCAGTCAGGC TTTCTTTCAC	3780
191 192	AGATGTGGAT TGGCGATAAA AAACAACTGC TGACGCCGCT GCGCGATCAG TTCACCCGTG	3840
193 194	CACCGCTGGA TAACGACATT GGCGTAAGTG AAGCGACCCG CATTGACCCT AACGCCTGGG	3900
195 196	TCGAACGCTG GAAGGCGGCG GGCCATTACC AGGCCGAAGC AGCGTTGTTG CAGTGCACGG	3960
197 198	CAGATACACT TGCTGATGCG GTGCTGATTA CGACCGCTCA CGCGTGGCAG CATCAGGGGA	4020
199 200	AAACCTTATT TATCAGCCGG AAAACCTACC GGATTGATGG TAGTGGTCAA ATGGCGATTA	4080
	CCGTTGATGT TGAAGTGGCG AGCGATACAC CGCATCCGGC GCGGATTGGC CTGAACTGCC	4140
203 204	AGCTGGCGCA GGTAGCAGAG CGGGTAAACT GGCTCGGATT AGGGCCGCAA GAAAACTATC	4200
205	CCGACCGCCT TACTGCCGCC TGTTTTGACC GCTGGGATCT GCCATTGTCA GACATGTATA	4260

206

RAW SEQUENCE LISTING PATENT APPLICATION US/09/242,977

DATE: 02/04/2000 TIME: 02:02:37

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200							
207 208	CCCCGTACGT	CTTCCCGAGO	GAAAACGGT	C TGCGCTGCGG	GACGCGCGA	A TTGAATTATG	4320
209 210	GCCCACACCA	A GTGGCGCGGC	GACTTCCAG	TCAACATCAG	CCGCTACAGT	CAACAGCAAC	4380
211 212	TGATGGAAAC	CAGCCATCGC	CATCTGCTGC	C ACGCGGAAGA	AGGCACATGG	G CTGAATATCG	4440
213	ACGGTTTCCA	TATGGGGATT	' GGTGGCGAC	ACTCCTGGAG	CCCGTCAGTA	TCGGCGGAAT	4500
214 215	TACAGCTGAG	CGCCGGTCGC	TACCATTACC	AGTTGGTCTG	GTGTCAAAAA	TAATAATAAC	4560
216 217	CGGGCAGGCC	ATGTCTGCCC	GTATTTCGCG	TAAGGAAATC	CATTATGTAC	TATTTAAAAA	4620
218 219	ACACAAACTT	' TTGGATGTTC	GGTTTATTCT	TTTTCTTTTA	CTTTTTTATC	ATGGGAGCCT	4680
220 221	ACTTCCCGTT	TTTCCCGATT	TGGCTACATG	ACATCAACCA	TATCAGCAAA	AGTGATACGG	4740
222 223	GTATTATTT	TGCCGCTATT	TCTCTGTTCT	' CGCTATTATT	CCAACCGCTG	TTTGGTCTGC	4800
224 225						GATAAGATAC	4860
226 227						TATTTGTGAA	4920
228 229						AGTTAACAAC	4980
230 231						TTTTTCGGAT	
232 233						AGGAACCCCT	5040
234							5100
235 236	AGTGATGGAG	TTGGCCACTC	CCTCTCTGCG	CGCTCGCTCG	CTCACTGAGG	CCGGGCGACC	5160
237 238	AAAGGTCGCC	CGACGCCCGG	GCTTTGCCCG	GGCGGCCTCA	GTGAGCGAGC	GAGCGCGCAG	5220
239	CAGATCTGGA	AGGTGCTGAG	GTACGATGAG	ACCCGCACCA	GGTGCAGACC	CTGCGAGTGT	5280
240 241	GGCGGTAAAC	ATATTAGGAA	CCAGCCTGTG	ATGCTGGATG	TGACCGAGGA	GCTGAGGCCC	5340
242 243	GATCACTTGG	TGCTGGCCTG	CACCCGCGCT	GAGTTTGGCT	CTAGCGATGA	AGATACAGAT	5400
244 245	TGAGGTACTG	AAATGTGTGG	GCGTGGCTTA	AGGGTGGGAA	AGAATATATA	AGGTGGGGGT	5460
246 247						CAACTCGTTT	5520
248 249						CGGGGTGCGT	5580
250 251						CTCTACTACC	5640
252 253						CGCCGCTTCA	
254							5700
255 256				ACTGACTTTG			5760
257 258	AGCAGTGCAG	CTTCCCGTTC	ATCCGCCCGC	GATGACAAGT	TGACGGCTCT	TTTGGCACAA	5820

SEQUENCE VERIFICATION REPORT PATENT APPLICATION US/09/242,977

DATE: 02/04/2000 TIME: 02:02:38

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Original Text

28

Wrong application Serial Number

(A) APPLICATION NUMBER: US